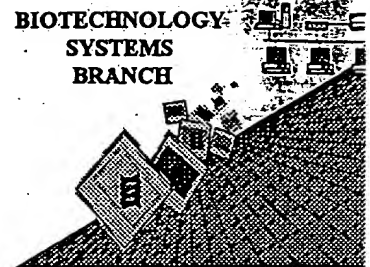


14

RAW SEQUENCE LISTING ERROR REPORT



p#5

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/407 804

Art Unit / Team No. :

1653

Date Processed by STIC:

5/15/2000

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MAY 30 2000

TECH CENTER 1600/2000

THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.

PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,**
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY**

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

1653

RAW SEQUENCE LISTING DATE: 05/15/2000
PATENT APPLICATION: US/09/407,804 TIME: 12:06:04

Input Set : A:\241 190-us -- sequence listing.txt
Output Set: N:\CRF3\05152000\I407804.raw

Does Not Comply
Corrected Diskette Needed

4 <110> APPLICANT: Jerry Pelletier
5 Philippe Gros
6 Michael DuBow
8 <120> TITLE OF INVENTION: DNA SEQUENCES FROM STAPHYLOCOCCUS AUREUS
9 BACTERIOPHAGE 77 THAT ENCODE ANTI-MICROBIAL POLYPEPTIDES
12 <130> FILE REFERENCE: 241/190US
14 <140> CURRENT APPLICATION NUMBER: 09/407,804
15 <141> CURRENT FILING DATE: 1999-09-28
17 <150> PRIOR APPLICATION NUMBER: 60/110,992
18 <151> PRIOR FILING DATE: 1998-12-03
E--> 20 <160> NUMBER OF SEQ ID NOS: 30
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

see next page - 29 shown in file

ERRORED SEQUENCES

09/407,804

Last sequence in file

<210> 29
<211> 237
<212> PRT
<213> Staphylococcus aureus bacteriophage 77

<400> 29

Met	Thr	His	Asn	Ile	Glu	Lys	Arg	Ile	Asn	Lys	Leu	Lys	Thr	Ser	Gly
1			5					10						15	
Asn	Pro	Lys	Phe	Lys	Lys	Leu	Asp	Ser	Asp	Ile	His	Tyr	Leu	Leu	Lys
		20					25						30		
Arg	Phe	Glu	Gly	Glu	Lys	Asn	His	Lys	Gly	Phe	Tyr	Pro	Lys	Phe	Lys
		35				40						45			
Gln	Gly	Glu	Ile	Val	Phe	Val	Asp	Phe	Gly	Ile	Asn	Val	Asn	Lys	Glu
		50				55					60				
Phe	Ser	Asn	Ser	His	Phe	Ala	Ile	Val	Met	Asn	Lys	Asn	Asp	Ser	Asn
65					70				75						80
Thr	Glu	Asp	Ile	Val	Asn	Val	Ile	Pro	Leu	Ser	Ser	Lys	Glu	Asn	Lys
			85					90						95	
Lys	Tyr	Leu	Lys	Met	Asn	Phe	Asp	Leu	Lys	Trp	Glu	Tyr	Tyr	Leu	Arg
			100					105						110	
Leu	Phe	Leu	Asn	Leu	Ile	Ser	Ala	Gln	Asn	Asn	Ser	Ala	Ile	Leu	Lys
			115				120						125		
Glu	Val	Phe	Asp	Lys	Lys	Tyr	Gln	Lys	Asn	Asn	Thr	Glu	Phe	Ile	Thr
			130				135					140			
Lys	Asp	Tyr	Phe	Ile	Glu	Phe	Ile	Ser	Asp	Ser	Leu	Glu	Ile	Glu	Asn
145					150				155						160
Lys	Leu	Asn	Lys	Ile	Asp	Arg	Asn	Ile	Asn	Asn	Ile	Val	Ser	Ala	Ile
				165					170						175
Asp	Lys	Val	Lys	Lys	Leu	Lys	Gly	Asn	Ser	Tyr	Ala	Cys	Ile	Asn	Ser
			180					185					190		
Phe	Gln	Pro	Ile	Ser	Lys	Phe	Arg	Ile	Arg	Lys	Val	Leu	Pro	Gln	Lys
			195				200					205			
Ile	Lys	Asn	Pro	Val	Ile	Asp	Ser	Ser	Asp	Ile	Met	Leu	Leu	Ile	Asn
		210				215					220				
Arg	Ile	Asn	Asn	Asn	Ile	Leu	Gln	Ile	Pro	Asp	Ile	Arg			
225					230						235				

RECEIVED

MAY 30 2003

TECH CENTER 1600/2900

VERIFICATION SUMMARY DATE: 05/15/2000
PATENT APPLICATION: US/09/407,804 TIME: 12:06:05

Input Set : A:\241 190-us -- sequence listing.txt
Output Set: N:\CRF3\05152000\I407804.raw

L:851 M:201 W: Mandatory field data missing, ORGANISM
L:852 M:280 W: Numeric Identifier already exists, Organism not replaced.
L:20 M:203 E: (20) Calc# of Seq. differs from actual, <160> Number Of Sequences:Input (30) Counted (29)